

SEQUENCE LISTING

SEQ ID NO:1

human CNG3B amino acid sequence

5 MFKSLTKVNVKVKPIGENNENEQSSRRNEEGSHPSNQSQQTTAQEENKGEEKSLKTKSTPVTSE
 EEPHTNIQDKLSKKNSSGDLTTNPDPQNAAEPTGTVPEQKEMDPGKEGPNSPQNKPPAAPVI
 NEYADAQLHNLVKRMRQRTALYKKKLVEGDLSSPEASPQTAKPTAVPPVKESDDKPTHEYYR
 LLWFKVKKMPLTEYLKRIKLPNSIDSYTDRLYLLWLLVTLAYNWNCFIPLRLVFPYQTAD
 10 NIHYWLIADIICDIIYLYDMLFIQPRLQFVRGGDIIVDSNELRKHYRTSTKFQLDVASIIPF
 DICYLFFGFNPMFRANRMLKYTSFFEFNHLESIMDKAYIYRVIRTTGYLLFILHINACVYY
 WASNYEGIGTTRWVYDGEENEYLRICYWAVRTLITIGGLPEPQTLFEIVFQLLNFFSGVFVF
 SSLIGQMRDVI GAATANQNYFRACMDDTIAYMNYSIPKLVQKRVRTWYEYTWDSQRMIDES
 DLLKTLPTTVQLALADVNFSIIISKVDLFKGCQTQMIYDMLRLKSVLYLPGDFVCKKGEIG
 KEMYIIKHGEVQVLGGPDGTVLVTLKAGSVFGEISLLAAGGGNRRTANVVAHGAFANLLTLD
 15 KKTLEILVHYPDSEIRILMKARVLLKQAKTAEATPPRKDLALLFPPKEETPKLFKTLGG
 TGKASLARLLKLKREQAAQKKENSEGEEEEGKENEDKQKENEDKQKENEDKQKENEDKDKGR
 EPEEKPLDRPECTASPIAVEEEPHSVRRTVLPRTSRQSLIISMAPPAGEEVLTIIEVKEK
 AKQ

20

SEQ ID NO:2

complete human CNG3B nucleotide sequence

25 CATTTCTCTACCTTAAGGCACAGTCATAAATACAGAGGGTTTTTCAGAACCACCTCAGAGAAGATG
 TTTAAATCGCTGACAAAAGTCAACAAGGTGAAGCCTATAGGAGAGAACAATGAGAATGAACAAAG
 TTCTCGTCGGAATGAAGAAGGCTCTCACCCAAGTAATCAGTCTCAGCAAACCACAGCACAGGAAG
 AAAACAAAGGTGAAGAGAAATCTCTCAAACCAAGTCAACTCCAGTCACGTCTGAAGAGCCACAC
 ACCAACATACAAGACAAACTCTCCAAGAAAAATTCCTCTGGAGATCTGACCACAAACCTGACCC
 TCAAATGCAGCAGAACCAACTGGAACAGTGCCAGAGCAGAAGGAAATGGACCCCGGAAAGAAG
 30 GTCCAAACAGCCCAAAAACAAACCGCCTGCAGCTCCTGTTATAAATGAGTATGCCGATGCCAG
 CTACACAACCTGGTGAAAAGAAATGCGTCAAAGAACAGCCCTCTACAAGAAAAGTTGGTAGAGGG
 AGATCTCTCTCACCCGAAGCCAGCCCAAACTGCAAAGCCACGGCTGTACCACCAGTAAAG
 AAAGCGATGATAAGCCAACAGAACATTACTACAGGCTGTTGTGGTTCAAAGTCAAAAAGATGCCT
 TTAACAGAGTACTTAAAGCGAATTAAACTTCCAACAGCATAGATTCATACACAGATCGACTCTA
 35 TCTCCTGTGGCTCTTGCTTGCTCACTCTTGCTTATAACTGGAAGTCTGGTTTATACCACTGCGCC
 TCGTCTTCCCATATCAAACCGCAGACAACATACTACTGGCTTATTGCGGACATCATATGTGAT
 ATCATCTACCTTTATGATATGCTATTTATCCAGCCAGACTCCAGTTTGTAAGAGGAGGAGACAT
 AATAGTGGAATCAAATGAGCTAAGGAAACACTACAGGACTTCTACAAAATTTTCAGTTGGATGTGCG
 CATCAATAATACCATTGATATTTGCTACCTCTTCTTTGGGTTTAATCCAATGTTTAGAGCAAAT
 40 AGGATGTTAAAGTACACTTCATTTTTTTGAATTTAATCATCACCTAGAGTCTATAATGGACAAAGC
 ATATATCTACAGAGTTATTGCAACAACCTGGTACTTGTGTTTATTCTGCACATTAATGCCTGTG
 TTTATTACTGGGCTTCAAACCTATGAAGGAATTGGCACTACTAGATGGGTGTATGATGGGGAAGGA
 AACGAGTATCTGGATGTTATTATTGGGCAGTTGCAACTTTAATTACCATTGGTGGCCTTCCAGA
 ACCACAACTTTATTTGAAATTGTTTTTCAACTCTTGAATTTTTTTTCTGGAGTTTTTGTGTTCT
 45 CCAGTTTAATTGGTCAGATGAGAGATGTGATTGGAGCAGCTACAGCCAATCAGAACTACTTCCGC
 GCCTGCATGGATGACACCATTGCCTACATGAACAATTACTCCATTCTAAACTTGTGCAAAAGCG
 AGTTTCGACTTGGTATGAATATACATGGGACTCTCAAAGAATGCTAGATGAGTCTGATTTGCTTA
 AGACCCTACCAACTACGGTCCAGTTAGCCCTCGCCATTGATGTGAAGTTTCTAGCATCATCAGCAA
 GTCGACTTGTTCAAGGGTTGTGATACACAGATGATTTATGACATGTTGCTAAGATTGAAATCCGT

TCTCTATTTGCCTGGTGACTTTGTCTGCAAAAAGGGAGAAATTGGCAAGGAAATGTATATCATCA
 AGCATGGAGAAGTCCAAGTTCTTGGAGGCCCTGATGGTACTAAAGTTCTGGTTACTCTGAAAGCT
 GGGTCGGTGTTTGGAGAAATCAGCCTTCTAGCAGCAGGAGGAGGAAACCGTCGAACTGCCAATGT
 GGTGGCCACGGGTTTGCCAATCTTTTAACTCTAGACAAAAAGACCCTCCAAGAAATTCTAGTGC
 5 ATTATCCAGATTCTGAAAGGATCCTCATGAAGAAAGCCAGAGTGCTTTTAAAGCAGAAGGCTAAG
 ACCGCAGAAGCAACCCCTCCAAGAAAAGATCTTGCCCTCCTCTTCCCACCGAAAGAAGAGACACC
 CAAACTGTTTTAAACTCTCCTAGGAGGCACAGGAAAAGCAAGTCTTGCAAGACTACTCAAATTGA
 AGCGAGAGCAAGCAGCTCAGAAGAAAGAAAATTCTGAAGGAGGAGAGGAAGAAGGAAAAGAAAAT
 GAAGATAAACAAAAAGAAAATGAAGATAAACAAAAAGAAAATGAAGATAAAGGAAAAGAAAATGA
 10 AGATAAAGATAAAGGAAGAGAGCCAGAAGAGAAGCCACTGGACAGACCTGAATGTACAGCAAGTC
 CTATTGCAGTGGAGGAAGAACCCCACTCAGTTAGAAGGACAGTTTTTACCCAGAGGGACTTCTCGT
 CAATCACTCATTATCAGCATGGCTCCTTCTGCTGAGGGCGGAGAAGAGGTTCTTACTATTGAAGT
 CAAAGAAAAGGCTAAGCAATAAATGTTTGATTATCTTTAGATGTGATATAGCTAGTTCCCAAAGT
 GATTGTACCTAGGATTGTAACCTAAATTAACGAGGGGAAACGACATGCTGGGACCCTTGAGAAAC
 15 GAAAGGCAAATCCCTAGCTTAGTTTTCTAGGACTTATCTGAGAGTGTGATTTTCATGCAGTGGTAAT
 AAGAAGATTATTAAGCAAAAAA

SEQ ID NO:3

20 human CNG3B coding sequence

ATGTTTAAATCGCTGACAAAAGTCAACAAGGTGAAGCCTATAGGAGAGAACAATGAGAATGAACA
 AAGTTCTCGTCGGAATGAAGAAGGCTCTCACCCAAGTAATCAGTCTCAGCAAACCACAGCACAGG
 AAGAAAACAAAGGTGAAGAGAAATCTCTCAAACCAAGTCAACTCCAGTCACGTCTGAAGAGCCA
 25 CACACCAACATACAAGACAACTCTCCAAGAAAAATTCTCTGGAGATCTGACCACAAACCTTGA
 CCTCAAATGCAGCAGAACCAACTGGAACAGTGCCAGAGCAGAAGGAAATGGACCCCGGGAAAG
 AAGGTCCAAACAGCCCACAAAACAAACCGCCTGCAGCTCCTGTTATAAATGAGTATGCCGATGCC
 CAGCTACACAACCTGGTGAAAAGAATGCGTCAAAGAACAGCCCTCTACAAGAAAAAGTTGGTAGA
 GGGAGATCTCTCTCACCCGAAGCCAGCCACAACTGCAAAGCCACGGCTGTACCACCAGTAA
 30 AAGAAAGCGATGATAAGCCAACAGAACATTACTACAGGCTGTTGTGGTTCAAAGTCAAAAAGATG
 CCTTTAACAGAGTACTTAAAGCGAATTAACTTCCAAACAGCATAGATTCATACACAGATCGACT
 CTATCTCCTGTGGCTCTTGCTTGTCACTCTTGCTATAACTGGAAGTCTGTTTATACCACTGC
 GCCTCGTCTTCCCATATCAAACCGCAGACAACATACTACTGGCTTATTGCGGACATCATATGT
 GATATCATCTACCTTTATGATATGCTATTTATCCAGCCAGACTCCAGTTTGTAAGAGGAGGAGA
 35 CATAATAGTGGATTCAAATGAGCTAAGGAAACACTACAGGACTTCTACAAAATTTTCAGTTGGATG
 TCGCATCAATAATACCATTTGATATTGCTACCTCTTCTTTGGGTTTAATCCAATGTTTAGAGCA
 AATAGGATGTTAAAGTACACTTCATTTTTGAATTTAATCATCACCTAGAGTCTATAATGGACAA
 AGCATATATCTACAGAGTTATTGAACAACCTGGATACTTGCTGTTTATTCTGCACATTAATGCCT
 GTGTTTATTACTGGGCTTCAAAGCTATGAAGGAATTGGCACTACTAGATGGGTGTATGATGGGGAA
 40 GGAAACGAGTATCTGAGATGTTATTATTGGGCAGTTGCAACTTTAATTACCATTGGTGCCCTTCC
 AGAACCACAACTTTATTTGAAATTGTTTTCAACTCTTGAATTTTTTTCTGGAGTTTTTGTGT
 TCTCCAGTTTAATTGGTCAGATGAGAGATGTGATTGGAGCAGCTACAGCCAATCAGAACTACTTC
 CGCGCCTGCATGGATGACACCATTGCCTACATGAACAATTACTCCATTCTAAACTTGTGCAAAA
 GCGAGTTTCGACTTGGTATGAATATACATGGGACTCTCAAAGAATGCTAGATGAGTCTGATTGTC
 45 TTAAGACCCTACCAACTACGGTCCAGTTAGCCCTCGCCATTGATGTGAACTTCAGCATCATCAGC
 AAAGTCGACTTGTTCAGGGTTGTGATACACAGATGATTTATGACATGTTGCTAAGATTGAAATC
 CGTTCTCTATTTGCCTGGTGACTTTGTCTGCAAAAAGGGAGAAATTGGCAAGGAAATGTATATCA
 TCAAGCATGGAGAAGTCCAAGTTCTTGGAGGCCCTGATGGTACTAAAGTTCTGGTTACTCTGAAA
 GCTGGGTGGTGTTTGGAGAAATCAGCCTTCTAGCAGCAGGAGGAGGAAACCGTCGAACTGCCAA
 50 TGTGGTGGCCACGGGTTTGCCAATCTTTTAACTCTAGACAAAAAGACCCTCCAAGAAATCTAG
 TGCATTATCCAGATTCTGAAAGGATCCTCATGAAGAAAGCCAGAGTGCTTTTAAAGCAGAAGGCT

AAGACCGCAGAAGCAACCCCTCCAAGAAAAGATCTTGCCCTCCTCTTCCCACCGAAAGAAGAGAC
ACCCAAACTGTTTAAAACTCTCCTAGGAGGCACAGGAAAAGCAAGTCTTGCAAGACTACTCAAAT
TGAAGCGAGAGCAAGCAGCTCAGAAGAAAAGAAAATTCTGAAGGAGGAGAGGAAGAAGGAAAAGAA
AATGAAGATAAACA AAAAGAAAATGAAGATAAACAAAAGAAAATGAAGATAAAGGAAAAGAAAA
5 TGAAGATAAAGATAAAGGAAGAGAGCCAGAAGAGAAGCCACTGGACAGACCTGAATGTACAGCAA
GTCCTATTGTCAGTGGAGGAAGAACCCCACTCAGTTAGAAGGACAGTTTTTACCCAGAGGGACTTCT
CGTCAATCACTCATTATCAGCATGGCTCCTTCTGCTGAGGGCGGAGAAGAGGTTCTTACTATTGA
AGTCAAAGAAAAGGCTAAGCAATAA

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SEQ ID NO:4

Oligo 1 (sense strand primer)

TCTATCTCCTGTGGCTCTTGCTTGTC

15

SEQ ID NO:5

Oligo 2 (antisense strand primer)

GAGTCTGGGCTGGATAAATAGCATATC

20

SEQ ID NO:6

Oligo 3 (sense strand primer)

AGGAATTGGCACTACTAGATGGGTG

25

SEQ ID NO:7

30 Oligo 4 (antisense strand primer)

TTCATGAGGATCCTTTCAGAATCTGG

35 **SEQ ID NO:8**

Oligo 5 (sense strand primer)

GGAAACCGTCGAACTGCCAATGTGGT

5 **SEQ ID NO:9**

Oligo 6 (sense strand primer)

CGGGTTTGCCAATCTTTTAACTCTAGAC

10

SEQ ID NO:10

Oligo 7 (antisense strand primer)

GTCCGCAATAAGCCAGTAGTGTATG

15

SEQ ID NO:11

Oligo 8 (sense strand primer)

20 TGACAAGCTTCCGCCATGTTTAAATCGCTGACAAAAGTC

SEQ ID NO:12

Oligo 9 (antisense strand primer)

25

TGACGAATTCTCCCAGCATGTCGTTTCCCCTCGTTAA